

file copy

Nucleotide

Search for

☐ 1: X83543. H.sapiens APXL mR...
[gi:790999]

ProbeSet, Related Sequences, OMIM, Protein, PubMed, Taxonomy,
UniSTS, LinkOut

LOCUS HSAPXL 7445 bp mRNA linear PRI 10-DEC-1995

DEFINITION H.sapiens APXL mRNA.

ACCESSION X83543

VERSION X83543.1 GI:790999

KEYWORDS APXL gene.

SOURCE human.

ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 7445)

AUTHORS Schiaffino, M.V., Bassi, M.T., Rugarli, E.I., Renieri, A., Galli, L. and Ballabio, A.

TITLE Cloning of a human homologue of the *Xenopus laevis* APX gene from the ocular albinism type 1 critical region

JOURNAL Hum. Mol. Genet. 4 (3), 373-382 (1995)

MEDLINE 95315933

REFERENCE 2 (bases 1 to 7445)

AUTHORS Schiaffino, V.M.

TITLE Direct Submission

JOURNAL Submitted (19-DEC-1994) V.M. Schiaffino, T.I.G.E.M., via Olgettina 58, 20132 Milano, ITALY

FEATURES

source Location/Qualifiers

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Revised: October 24, 2001.

Disclaimer | Write to the Help Desk
NCBI | NLM | NIH

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 Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.
 1 (bases 1 to 156375)
 Tsai,S.F.
 Direct Submission
 Submitted (28-AUG-2000) Shih-Feng Tsai, National Yang-Ming
 University, Institute of Genetics, 155 Li-Rong St. Section 2,
 Peitou, Taipei, Taiwan 11221, Republic of China
 (E-mail:ymptsai@ym.edu.tw, URL:http://genome.ym.edu.tw/,
 Tel:886-2-28267043, Fax:886-2-28264930)
 These sequences are draft human sequences, not finished sequences.
 These sequences are unordered pieces. And gaps between the contigs
 of the same clone are represented as 100 N.
 * NOTE: This is a 'working draft' sequence. It currently
 * consists of 15 contigs. The true order of the pieces
 * is not known and their order in this sequence record is
 * arbitrary. Gaps between the contigs are represented as
 * runs of N, but the exact sizes of the gaps are unknown.
 * This record will be updated with the finished sequence
 * as soon as it is available and the accession number will
 * be preserved.

1 21185: contig of 21185 bp in length
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FEATURES
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Query Match 48.1%; Score 157.4; DB 2; Length 156375;
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RESULT 7
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 1 (bases 1 to 7445)
 Schiaffino,M.V., Bassi,M.T., Ruggeri,E.I., Renneri,A., Galli,L. and
 Balibio,A.
 Cloning of a human homologue of the Xenopus laevis APX gene from
 the ocular albinism type I critical region
 Hum. Mol. Genet. 4 (3), 373-382 (1995)
 95315933
 2 (bases 1 to 7445)
 Schiaffino,V.M.
 Direct Submission
 Submitted (19-DEC-1994) V.M. Schiaffino, T.I.G.E.M., via Olgettina
 58, 20132 Milano, ITALY
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